

Questionnaire

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Your institution: Universidad Nacional de Educación a Distancia (UNED)

Your position: Profesor tutor

Undergraduate student	<input type="checkbox"/>
PhD student	<input type="checkbox"/>
Technician	<input type="checkbox"/>
Postdoc	<input type="checkbox"/>
Faculty member	<input checked="" type="checkbox"/>

Question 1: In a scale from 1 - 10, how important do you think it is to have quick access to the following type of information about each gene or protein?

- a. Protein-protein interactions [1-10]: 8
- b. General function of the gene/protein [1-10]: 8
- c. Diseases a gene/protein is involved in [1-10]: 9
- d. Biochemical pathways a gene/protein is part of [1-10]: 8
- e. Cell types/tissues where your gene/protein is expressed [1-10]: 7
- f. Your protein's 3D structure (PDB) [1-10]: 5
- g. Popularity of the gene/protein in **social networks (Twitter, Facebook)** [1-10]: 4
- h. Knowing the **average impact factor** of the journals where a particular gene/protein is normally published [1-10]: 6
- i. The **relative scientific weight (e.g. by h-index)** of the scientists that work on your gene/protein [1-10]: 6
- j. How popular your gene/protein is in recently awarded grants (this is public information once a grant is awarded) [1-10]: 7
- k. What other genes/proteins are discussed in the context of your protein [1-10]: 8

- l. How your gene/protein is regulated at the transcriptional level [1-10]: 8
- m. How your gene/protein is regulated post-translationally (phosphorylation, ubiquitination) [1-10]: 7
- n. What is the **most popular type of experiment** other scientists typically do on your gene/protein [1-10]: 8
- o. What **biochemical kits** are available for doing these experiments [1-10]:7
- p. Other (explain what type of information) [1-10]:

Question 2: What websites do you visit the most when analysing your list of genes/proteins?

What type of information do you expect to get from each of these websites?

Some examples of websites include:

- Ensembl (www.ensembl.org)
- NCBI's Entrez (<https://www.ncbi.nlm.nih.gov/Class/MLACourse/Original8Hour/Entrez/>)
- NCBI Databases (<https://www.ncbi.nlm.nih.gov/search/>)
- EuPathDB.org (Eukaryotic Pathogens Database)
- Galaxy (<https://usegalaxy.org/>)
- PubMed (<https://www.ncbi.nlm.nih.gov/pubmed/>)
- UniProt (<http://www.uniprot.org/>)
- KEGG Pathway Database (<https://www.genome.jp/kegg/pathway.html>)
- Any other resource you use routinely.

Website	Type of information sought	Priority in your analytical pipeline
PubMed	Find out what's been published about my gene or protein	1

Genome Browser Gateway https://genome-euro.ucsc.edu/	Transcriptional regulation	2
KEGG Pathway Database	Molecular interactions	3
<i>(expand the table as needed)</i>		

Question 3: How often do you perform these exploratory analyses on your genes or proteins:

- [a] Daily
- [b] Weekly
- [c] Monthly
- [d] Several times a year
- [e] Other (explain)

Question 4: If you could obtain the same type of information that you seek by doing these analyses in **5 minutes only**, how often would you now perform these analyses?

- [a] Daily
- [b] Weekly
- [c] Monthly
- [d] Several times a year
- [e] Other (explain)

Question 5: When you get a **list of genes/proteins** from a proteomics or a differential expression experiment, what **steps and tools** do you follow for the analysis?

1. Check biggest expression differences and work only on a couple of genes/proteins
2. Pubmed search
3. Transcriptional regulation (Genome Browser Gateway)
4. Molecular interactions (KEGG)
5. siRNAs and/or molecular inhibitors *in vitro*

Question 6: Now and related to the previous question, instead of telling me what steps you follow in your analysis, if I asked you **what type of information you would like to know** about each one of your genes or proteins, what would you be interested in knowing?

Here you can include some information you would love to have **but do not know** how to obtain it.

1. Main transcriptional regulators
2. Molecular interactions
3. Biochemical pathways
4. Related diseases
- 5.

Question 7: If you have a long list of genes/proteins from a high-throughput experiment you ran in the lab, **what are the most useful factors in determining the next follow-up experiment?**

E.g. how easy the potential experiment is, how relevant the cell type, how much money this would cost, etc.

1. How much money this would cost
2. How long this would take
3. How easy the potential experiment is
4. Colleagues that are experts in the experiments
- 5.

Question 8: What do you think is the **most competitive advantage against other competing laboratories** when trying to decide what the next follow-up experiment would be?

1. Money available to spend / people dedicated to that line of research

2. Relevance of the gene(s) / protein(s) selected

3.

4.

5.

__END OF QUESTIONNAIRE__